

Fig. 1

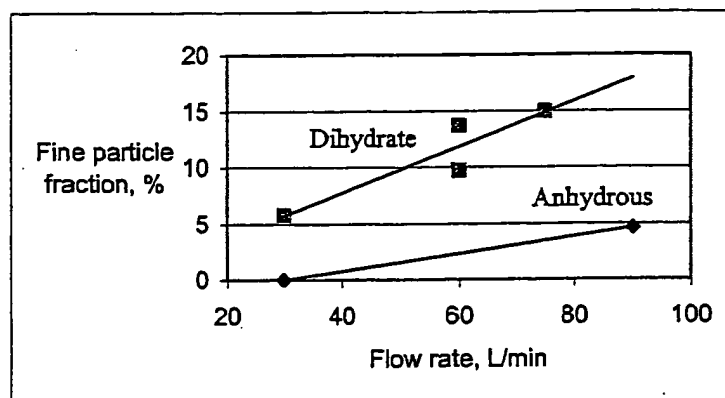


Fig. 2

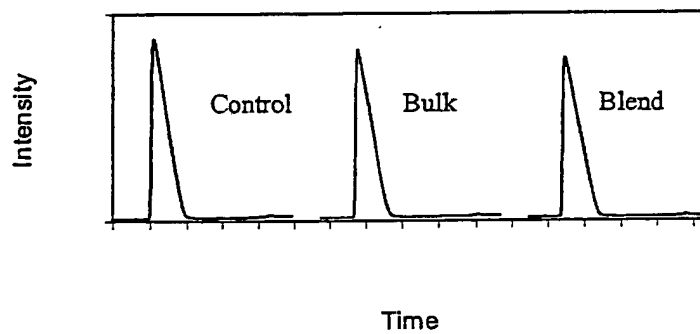


Fig. 3

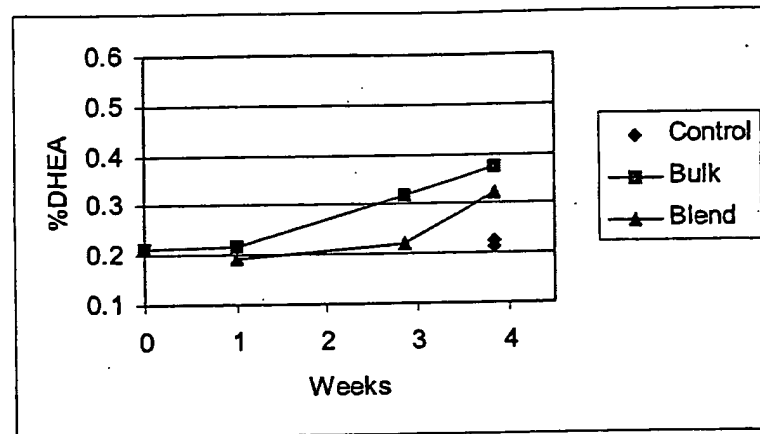


Fig. 4

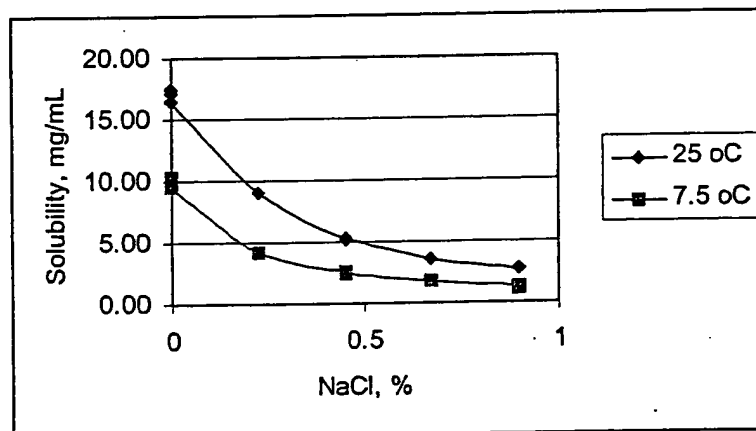


Fig. 5

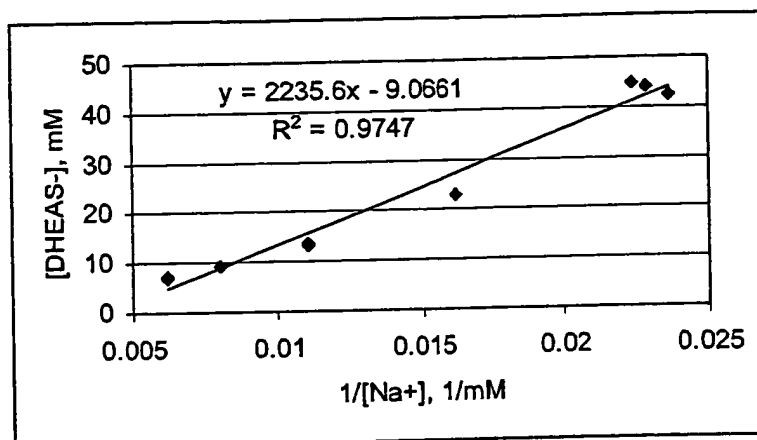


Fig. 6

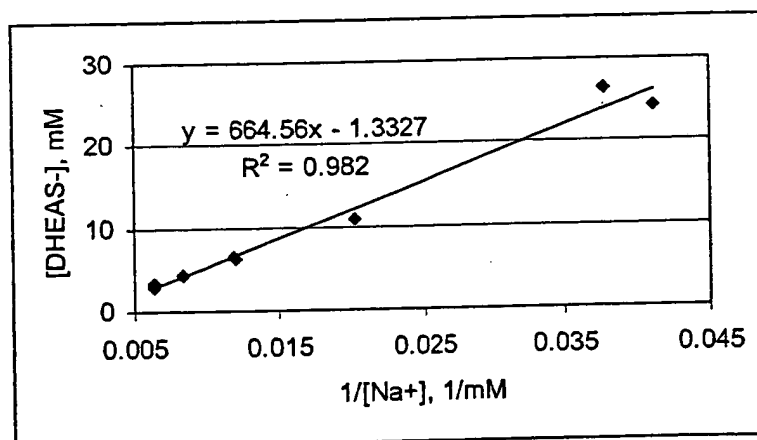


Fig. 7

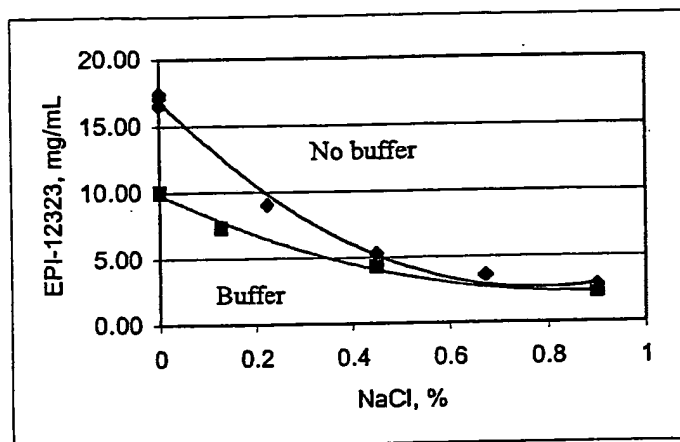


Fig. 8

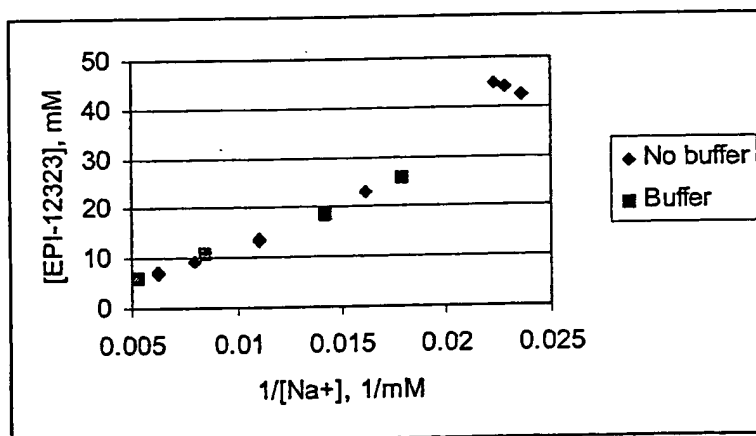


Fig. 9

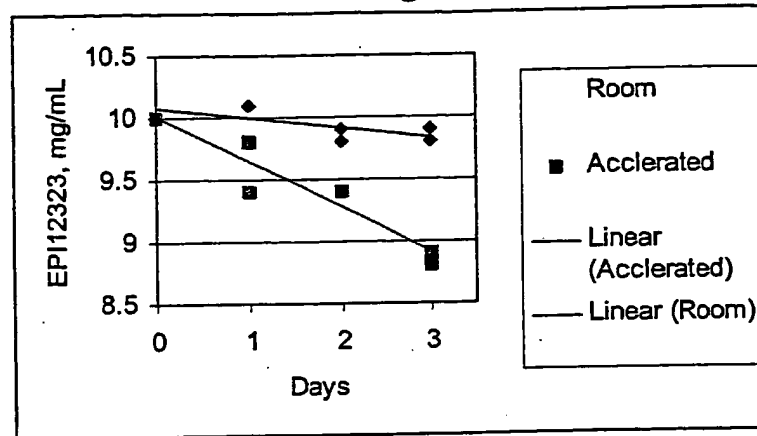


Fig. 10

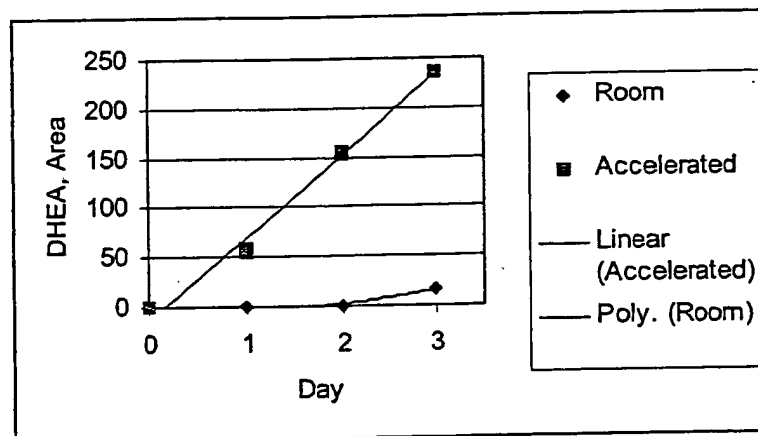


Fig. 11

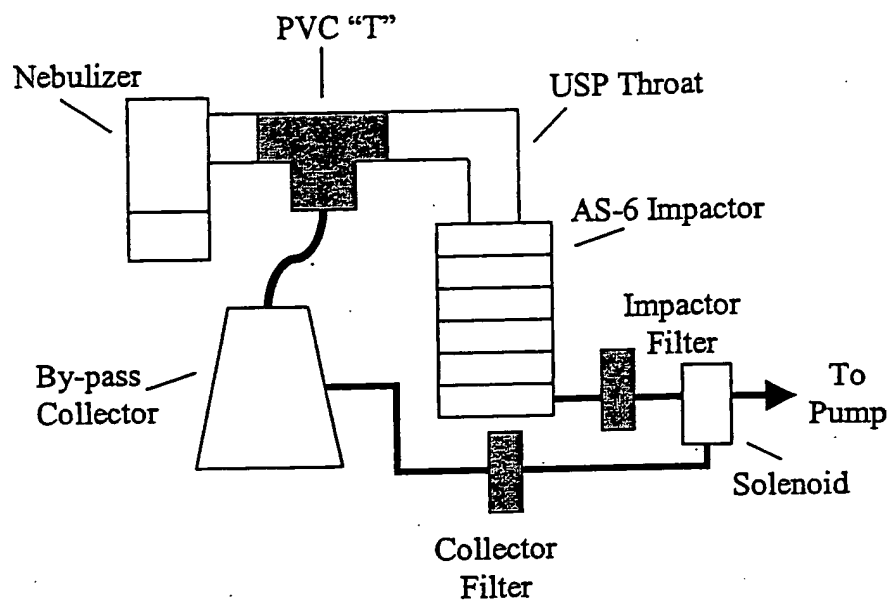


Fig. 12

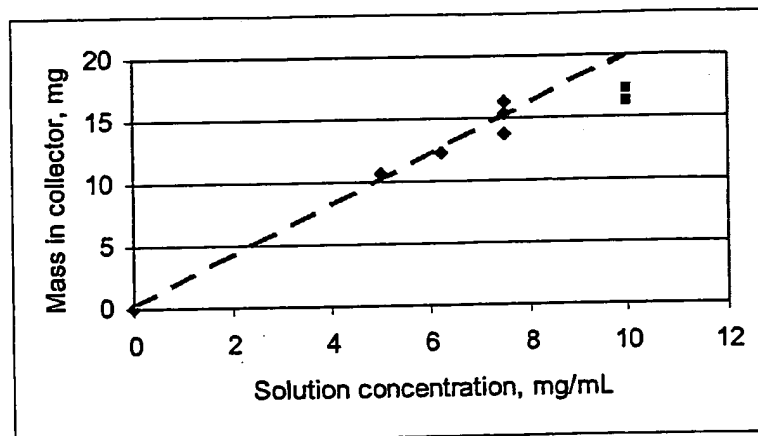


Fig. 13

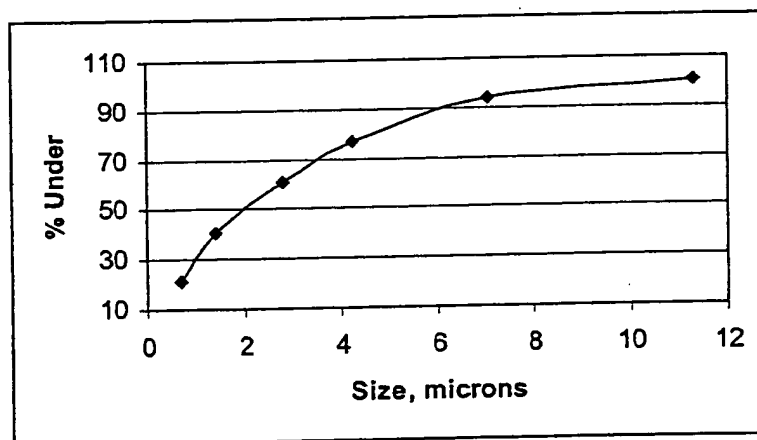


Fig. 14

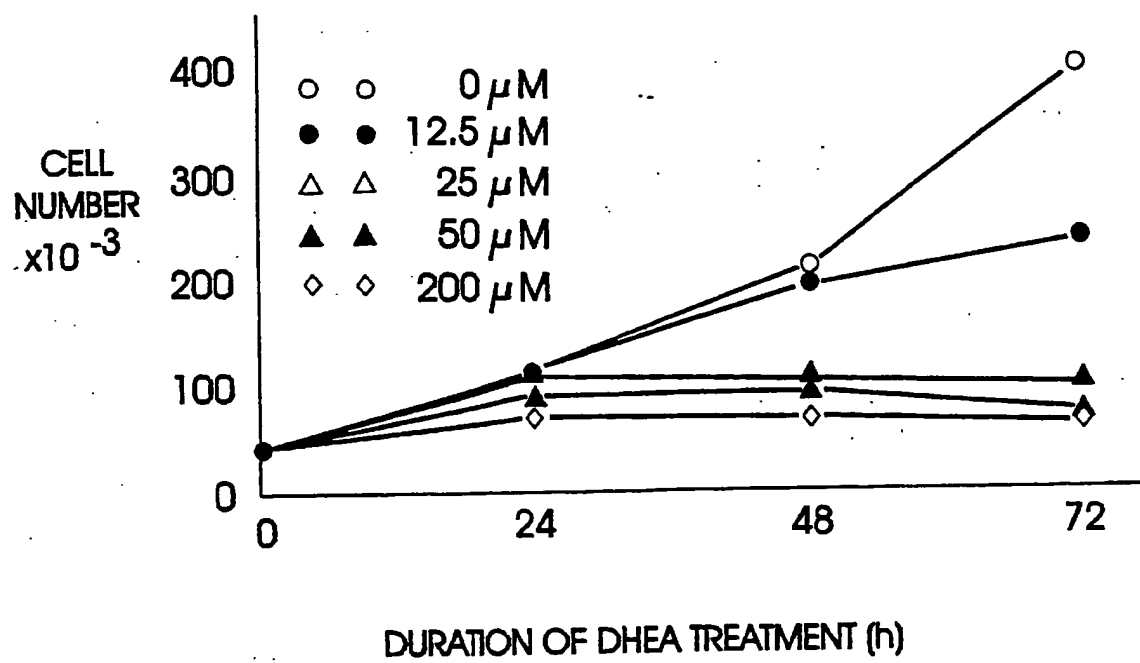


Fig. 15A

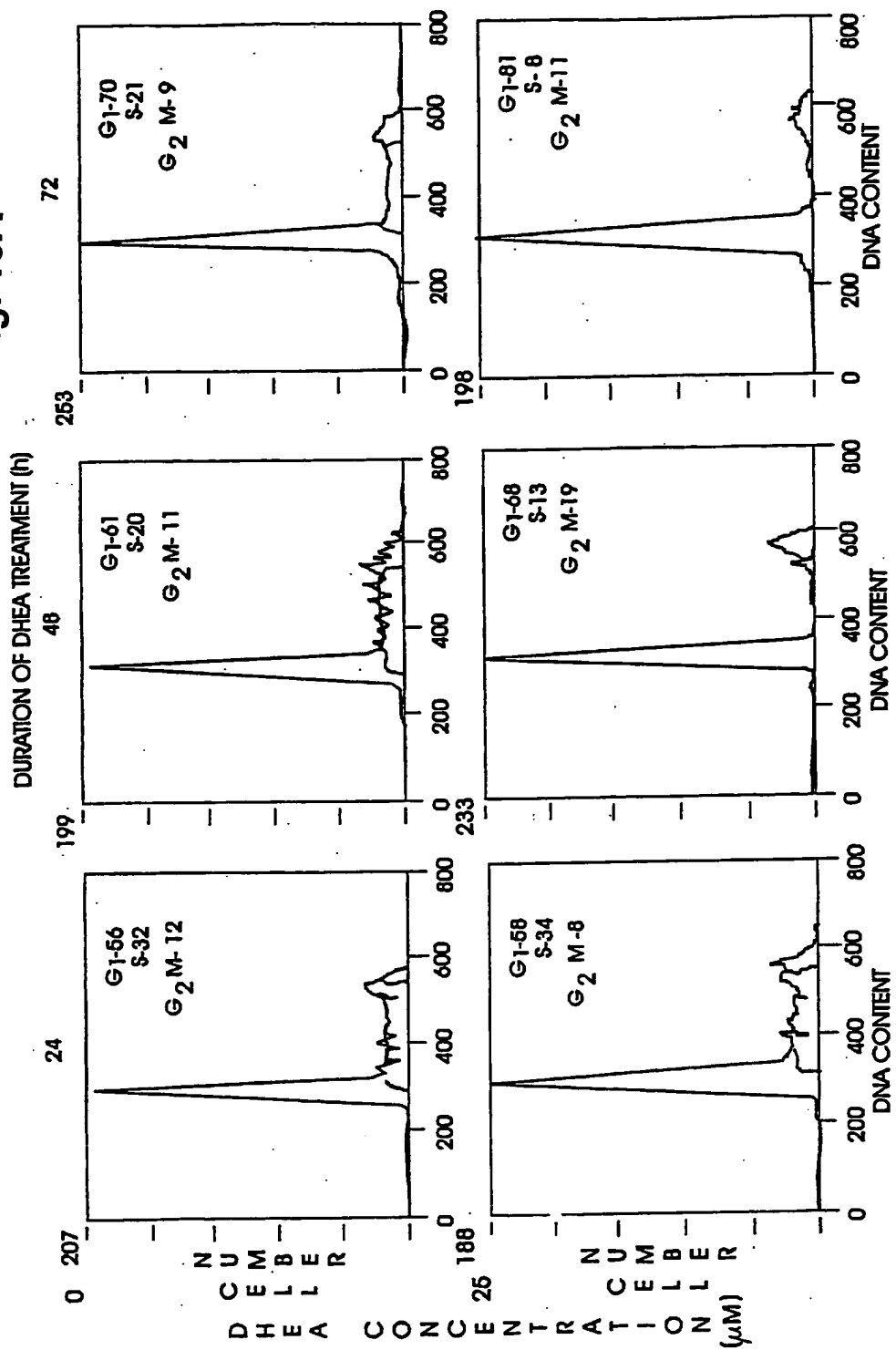
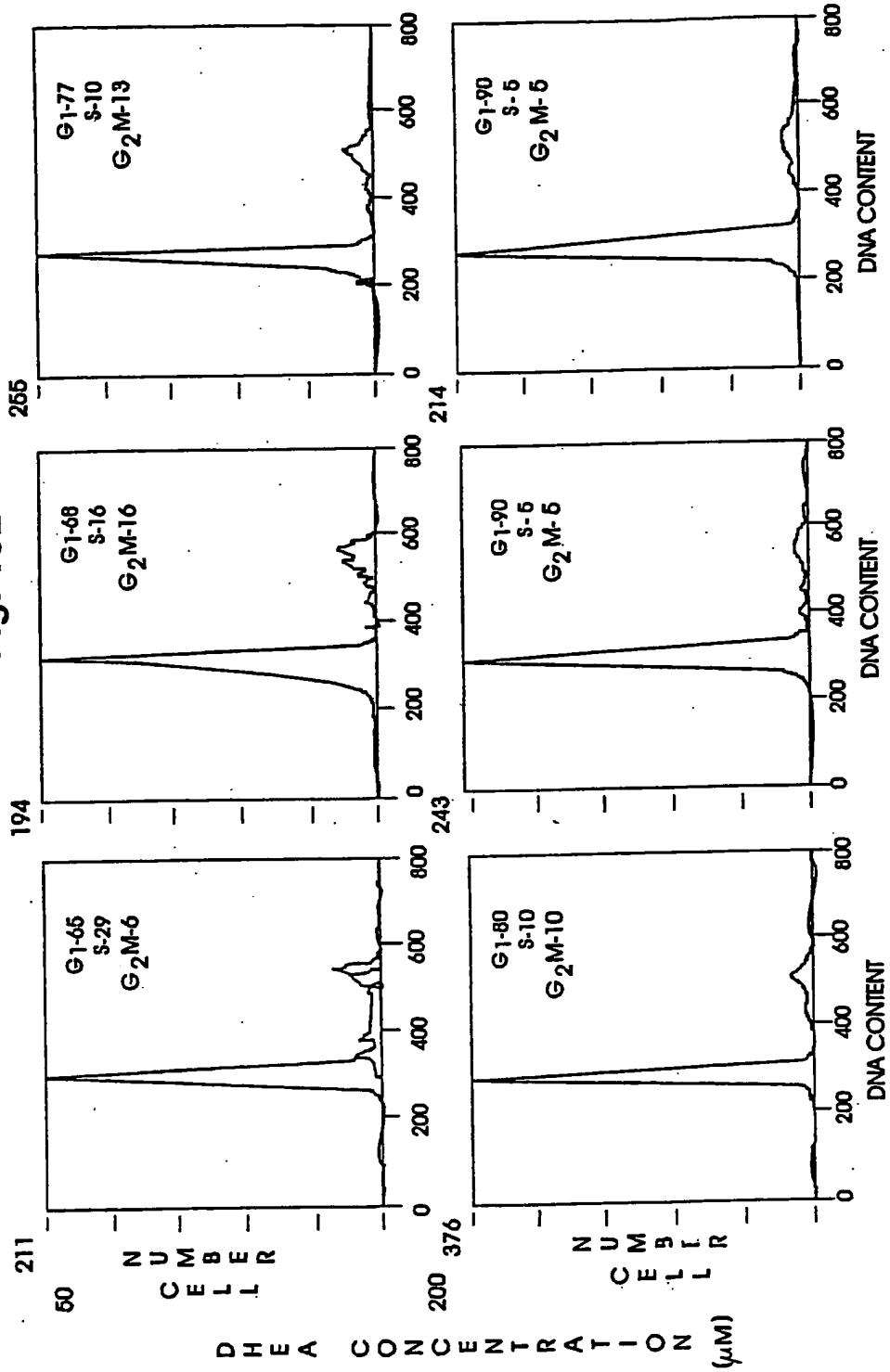


Fig. 15B



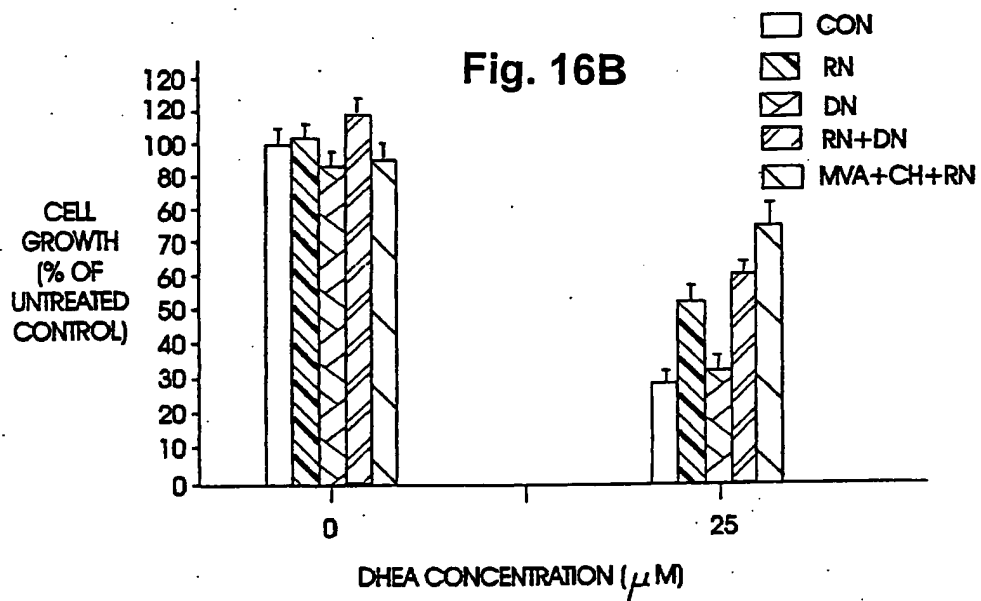
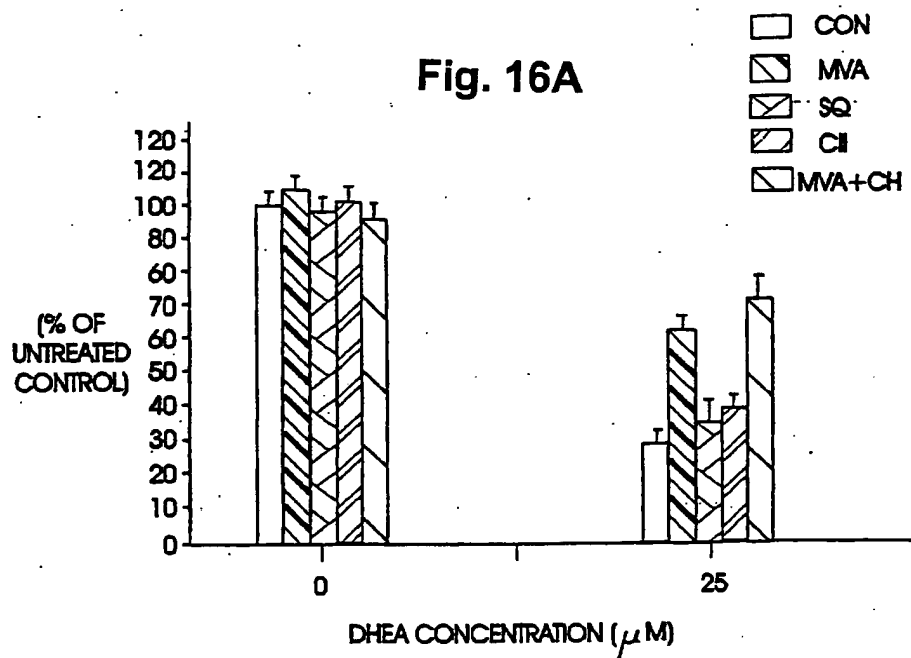


Fig. 17A

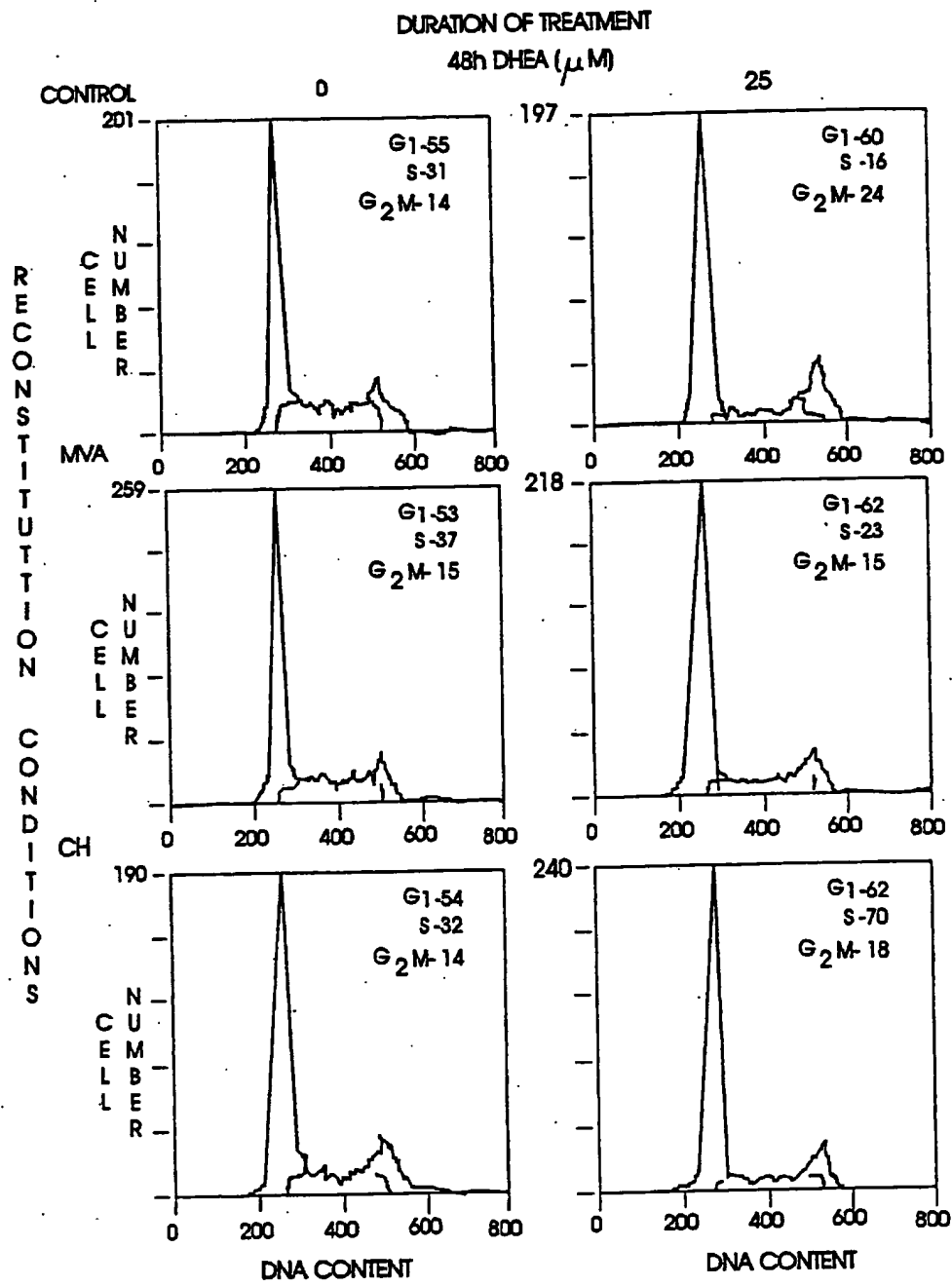
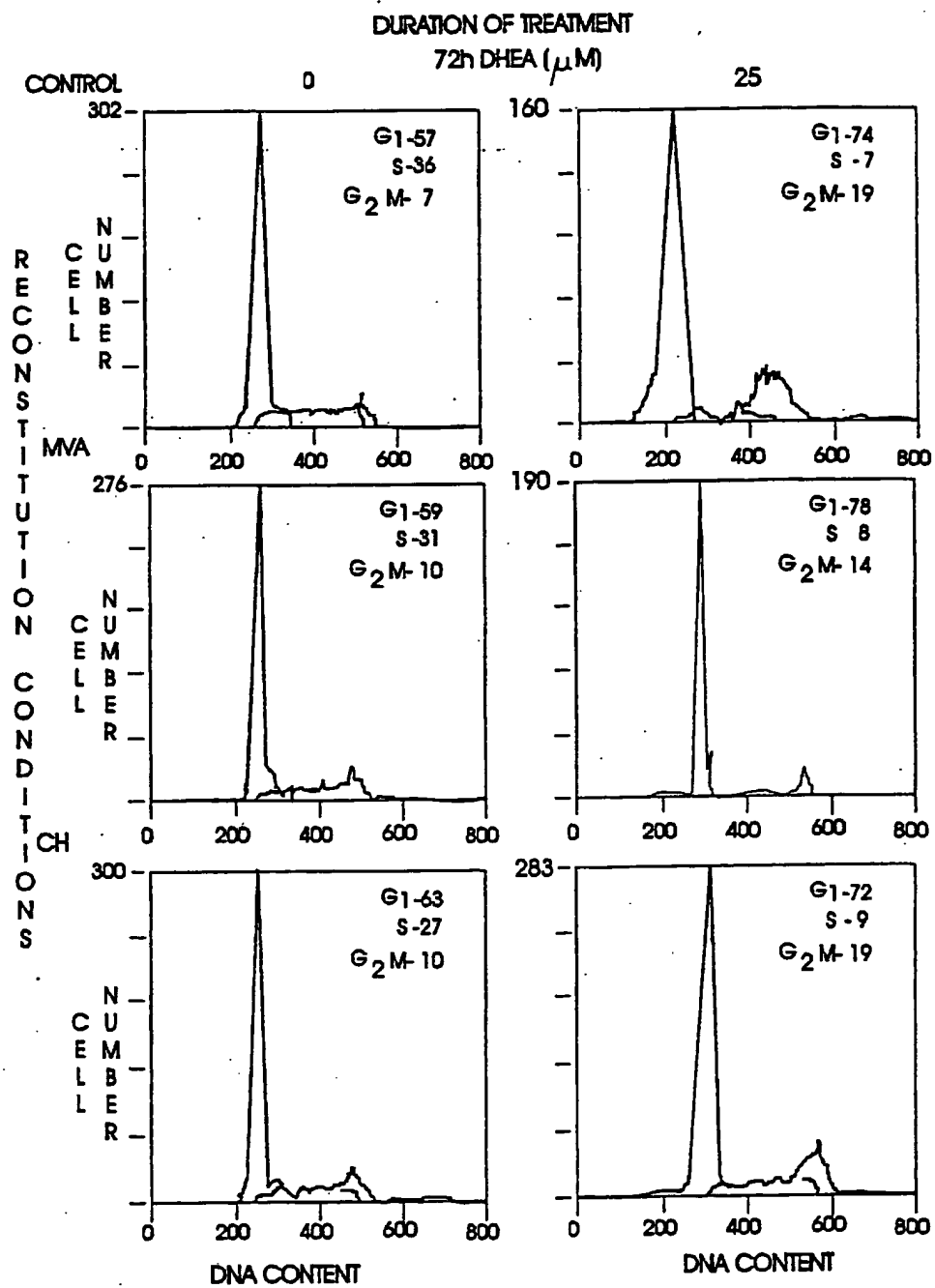


Fig. 17B



DURATION OF TREATMENT

The figure consists of six histograms arranged in a 3x2 grid. Each histogram plots 'CELL NUMBER' on the y-axis against 'DNA CONTENT' on the x-axis (ranging from 0 to 800). The histograms show the distribution of DNA content for different cell lines and conditions. The cell lines and conditions are labeled in each plot:

- Top Left:** G1-55, S-31, G₂M-14. Y-axis scale: 0 to 159. Condition: RN.
- Top Right:** G1-65, S-19, G₂M-16. Y-axis scale: 0 to 183. Condition: 25.
- Middle Left:** G1-53, S-32, G₂M-15. Y-axis scale: 0 to 165. Condition: MVA+CH.
- Middle Right:** G1-67, S-23, G₂M-10. Y-axis scale: 0 to 157. Condition: 25.
- Bottom Left:** G1-49, S-34, G₂M-17. Y-axis scale: 0 to 148. Condition: MVA+CH+RN.
- Bottom Right:** G1-64, S-28, G₂M-8. Y-axis scale: 0 to 183. Condition: 25.

Each histogram shows a primary peak at approximately 250 DNA content units and a secondary, broader peak around 500 DNA content units. The y-axis label 'CELL NUMBER' is oriented vertically on the left side of each plot. The x-axis label 'DNA CONTENT' is located at the bottom of the figure.

Fig. 17D

